

# Resource Summary Report

Generated by T1D on Aug 23, 2026

## Xenium Ranger

RRID:SCR\_028191

Type: Tool

### Proper Citation

Xenium Ranger (RRID:SCR\_028191)

### Resource Information

**URL:** <https://www.10xgenomics.com/support/software/xenium-ranger/latest>

**Proper Citation:** Xenium Ranger (RRID:SCR\_028191)

**Description:** Software suite for analyzing and visualizing in situ gene expression data produced by Xenium Analyzer. Xenium Ranger provides flexible off-instrument reanalysis of Xenium In Situ data.

**Resource Type:** software resource

**Keywords:** analyzing and visualizing in situ gene expression data, data produced by Xenium Analyzer,

**Availability:** Restricted

**Resource Name:** Xenium Ranger

**Resource ID:** SCR\_028191

**Record Creation Time:** 20260327T054533+0000

**Record Last Update:** 20260815T183315+0000

### Ratings and Alerts

No rating or validation information has been found for Xenium Ranger.

No alerts have been found for Xenium Ranger.

### Data and Source Information

**Source:** [SciCrunch Registry](#)

## Usage and Citation Metrics

We found 1 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [T1D](#) .

Wang BS, et al. (2026) Single-cell multiomic approaches define a gradual, spatially regulated epigenetic and transcriptional transition from embryonic to adult neural stem cells. Stem cell reports, 21(7), 102967.